

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/572,882  
Source: IFWP  
Date Processed by STIC: 3/31/06

***ENTERED***



IFWP

RAW SEQUENCE LISTING DATE: 03/31/2006  
 PATENT APPLICATION: US/10/572,882 TIME: 12:19:12

Input Set : A:\PTO.KD.txt  
 Output Set: N:\CRF4\03292006\J572882.raw

3 <110> APPLICANT: Bayer HealthCare AG  
 4 Golz, Stefan  
 5 Broggemeier, Ulf  
 6 Geerts, Andreas  
 8 <120> TITLE OF INVENTION: Diagnostics and Therapeutics for Diseases  
 Associated with  
 9 G-Protein Coupled Receptor ADIPOR2 (ADIPOR2)  
 11 <130> FILE REFERENCE: LeA:36 902  
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/572,882  
 C--> 13 <141> CURRENT FILING DATE: 2006-03-20  
 13 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/010383  
 14 <151> PRIOR FILING DATE: 2004-09-16  
 16 <150> PRIOR APPLICATION NUMBER: EP03021898.6  
 17 <151> PRIOR FILING DATE: 2003-09-27  
 19 <160> NUMBER OF SEQ ID NOS: 5  
 21 <170> SOFTWARE: PatentIn version 3.3  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 3500  
 25 <212> TYPE: DNA  
 26 <213> ORGANISM: Homo sapiens  
 28 <400> SEQUENCE: 1  
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 31 agactggctc aaggataatg acttccttctt gcatggacac cggcctctta tgccttcttt 120  
 33 ccgggcctgt ttaagagca ttttcagaat acacacagaa acaggcaaca tttggacaca 180  
 35 tcttttaggt tggatccct tcctgtgcct gggatcttt tataatgtttc gcccaaataat 240  
 37 ctcccttggc gcccctctgc aagagaagg ggtctttgga ttattttct taggagccat 300  
 39 tctctgcctt tcttttcat ggctcttcca cacagtctac tgccactcag agggggctc 360  
 41 tcggctcttc tctaaactgg attactctgg tattgtctt ctgattatgg gaagtttgt 420  
 43 tccttggctt tattattctt tctactgtaa tccacaacat tgcttcatct acttgattgt 480  
 45 catctgtgtg ctggcatttgc cagccattat agtctccag tggacatgt ttgccacccc 540  
 47 tcagtatcgg ggagtaagag caggagtgtt tttggccttgc ggctgagtg gaatcattcc 600  
 49 taccttgcac tatgtcatct cggagggtt ccttaaggcc gcacccatag ggcagatagg 660  
 51 ctgggtatg ctatggcca gcctctacat cacaggagct gcctgtatg ctggccggat 720  
 53 cccgaacgc ttttccctg gcaaatgtga catctggttt cactctcatc agctgttca 780  
 55 tatctttgtg gttgctggag ctttgcattca cttccatggt gtctcaaacc tccaggagtt 840  
 57 tcgtttcatg atcgccgggg gctgcagtga agaggatgca ctgtatacc taccagtctc 900  
 59 caggactat gaccctaaac cagggcctgc ggcacttgcg ggcctccctg ctggctactg 960  
 61 atgccagtac cagaggagcc ccaaaacttt gacagctcg tggctttgt gacggcccag 1020  
 63 gggctctgatg tggatcatga ctgagaagag aaaaacaaaa ataaatcata cctcaaagga 1080  
 65 tggagtgcattt caattggggaaa aaaaaggagac atagccaaaa ccctggctta ttcttggat 1140  
 67 ctactgatttgc cggcatttgc aagacccttgc gcaaaacttgc ttctgatcca tatcatattt 1200  
 69 attttagaa gatggcggaaa cagtttagct ggtggtttt tcttctccct ttctctct 1260  
 71 ctatgacaat aatacaaacc aatttaagt aacatttata tccgataagg ggtggagtg 1320  
 73 tgattttaaa tgcttttttgc ggagaacaaa gaaattaatg taaataagat ttctaaactgt 1380

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79	taggatcagg	tgatagcccc	ggaatgtaca	gtgtcttgg	gcaccaagat	gccttctaaa	1560										
81	ggctgacata	ccttggaccc	taatggggca	gagagtata	ccctagccca	gtggtgacat	1620										
83	gaccactccc	tttgggaggc	ctgaggtaga	ggggagtggt	atgtttttc	tcagtggaa	1680										
85	cagcacatga	gtgggtgaca	ggatgttaga	taaaggctct	atttagggtg	tcattgtcat	1740										
87	ttgagagact	gacacactcc	tagcagctgg	taaagggtg	ctggaggcca	tggaggagct	1800										
89	ctagaaacat	tagcatggc	tgatctgatt	acttcctggc	atcccgctca	cctttatggg	1860										
91	aagtcttatt	agagggatgg	gacagtttc	cataccctt	ctgtggagct	ctggaacact	1920										
93	ctctaaattt	ccctcttatta	aaaatcactg	ccctaactat	acttcctcct	tgagggaaata	1980										
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97	acgtgtccag	gttctcctg	agctcatcta	catagattgg	tagacccttc	ctttggattt	2100										
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103	tcagcactca	ctgaacctca	ctttacaggg	ataagagtgg	tgtggcattt	taaataacaat	2280										
105	ggtatgttat	tgccagggag	ttaggtacaa	gacgatggct	catgtcacag	gcctacctga	2340										
107	tacggtgtca	gagaaagtgg	tggggaaagg	atctggttca	tggaattctg	atcttggccc	2400										
109	ataggtgaac	caccaaaaata	gtgctcgagt	cttaggttac	tgtcatcaaa	gacttggat	2460										
111	gactccatta	tatcctgggg	ttgtgggtat	tagaactaaa	tatggaggtc	ctgagcatgg	2520										
113	ggactgggt	cctcagtagg	tgtttgggaa	tatgggaagg	gtctcctatt	tattcaatag	2580										
115	agtttctca	gttattttcc	tccttcgccc	ttgcaatctc	cagaaaagg	tggatctag	2640										
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119	cactgtccta	ggtgcttagt	ggaccgagca	aaagactcag	tggatgaact	ggtgcagtgc	2760										
121	ctgacagaat	aaagaacagt	attaatccct	ttgagaaaagc	atagtcacgc	aggacagtgg	2820										
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129	aacatagcac	aaacccttag	gaaaaatgaa	attaacatca	ctgatgtgt	atccagtaaa	3060										
131	atcccttt	ttcgggtgt	tatgtggca	tgtgccatt	tctatgtgt	tgtctacgtg	3120										
133	cagctacta	ccaacagcct	catgtgcact	tgacctgaca	gtgctcgctg	agaactctca	3180										
135	ccaggttggc	gcctgaatgc	cttactctca	gcagtcagag	gcttgcttc	tctgtgcaga	3240										
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141	acaaagcaca	tttttggga	tcatagaagg	ttggggttcc	agaaaggcat	ctgtgtatg	3420										
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149	<211>	LENGTH:	258														
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159	Glu	Thr	Gly	Asn	Ile	Trp	Thr	His	Leu	Leu	Gly	Cys	Val	Phe	Phe	Leu	
160						20				25					30		
163	Cys	Leu	Gly	Ile	Phe	Tyr	Met	Phe	Arg	Pro	Asn	Ile	Ser	Phe	Val	Ala	
164						35			40					45			
167	Pro	Leu	Gln	Glu	Lys	Val	Val	Phe	Gly	Leu	Phe	Phe	Leu	Gly	Ala	Ile	
168						50			55					60			

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171 Leu Cys Leu Ser Phe Ser Trp Leu Phe His Thr Val Tyr Cys His Ser  
 172 65 70 75 80  
 175 Glu Gly Val Ser Arg Leu Phe Ser Lys Leu Asp Tyr Ser Gly Ile Ala  
 176 85 90 95  
 179 Leu Leu Ile Met Gly Ser Phe Val Pro Trp Leu Tyr Tyr Ser Phe Tyr  
 180 100 105 110  
 183 Cys Asn Pro Gln Pro Cys Phe Ile Tyr Leu Ile Val Ile Cys Val Leu  
 184 115 120 125  
 187 Gly Ile Ala Ala Ile Ile Val Ser Gln Trp Asp Met Phe Ala Thr Pro  
 188 130 135 140  
 191 Gln Tyr Arg Gly Val Arg Ala Gly Val Phe Leu Gly Leu Gly Leu Ser  
 192 145 150 155 160  
 195 Gly Ile Ile Pro Thr Leu His Tyr Val Ile Ser Glu Gly Phe Leu Lys  
 196 165 170 175  
 199 Ala Ala Thr Ile Gly Gln Ile Gly Trp Leu Met Leu Met Ala Ser Leu  
 200 180 185 190  
 203 Tyr Ile Thr Gly Ala Ala Leu Tyr Ala Ala Arg Ile Pro Glu Arg Phe  
 204 195 200 205  
 207 Phe Pro Gly Lys Cys Asp Ile Trp Phe His Ser His Gln Leu Phe His  
 208 210 215 220  
 211 Ile Phe Val Val Ala Gly Ala Phe Val His Phe His Gly Val Ser Asn  
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 224 <211> LENGTH: 20  
 225 <212> TYPE: DNA  
 226 <213> ORGANISM: Artificial  
 228 <220> FEATURE:  
 229 <223> OTHER INFORMATION: Primer1 (forward primer)  
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 235 <210> SEQ ID NO: 4  
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 240 <220> FEATURE:  
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 244 cagtgcattc tcttcactgc 20  
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 256 agtttcgttt catgatcggc ggg 23

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/31/2006  
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Input Set : A:\PTO.KD.txt  
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### Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#: 3, 4, 5

**VERIFICATION SUMMARY** DATE: 03/31/2006  
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Input Set : A:\PTO.KD.txt  
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L:13 M:270 C: Current Application Number differs, Replaced Current Application No  
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date